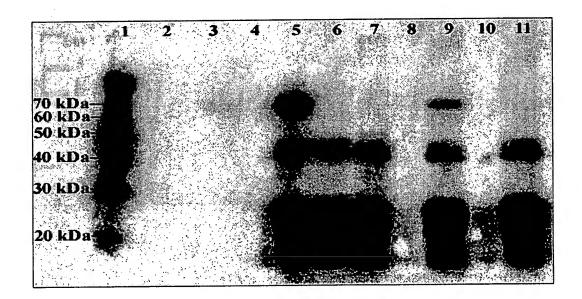
Fig. 1

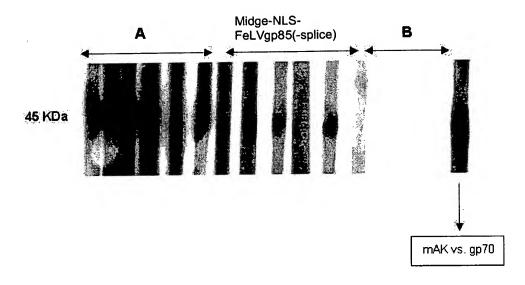


Fig. 2



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Fig. 3



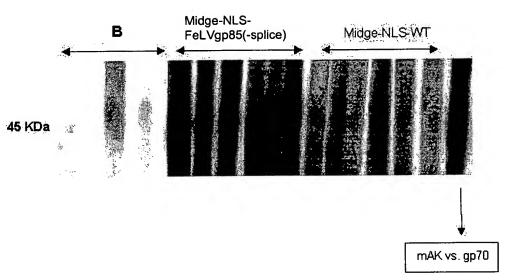


Fig. 4:	DNA sequence comparison of the wild type "gag" gene (Seq.ID2) against the codon-optimized "gag" gene (Seq.ID5).

codon-optimized gag gene (Seq.ibs).			
SeqID2 SeqID5	1	ATGGGCCAAACTATAACTACCCCCTTGAGCCTCACCCTCAACCACTGGTCTGAGGTTCAG	
SeqID2 SeqID5	61 61	GCACGGGCCCGTAATCAGGGTGTCGAAGTCCGGAAAAAGAAATGGATTACACTGTGTGAA ""CA""""A"G""C"""""C""G""G""GA""""G""""""""	
SeqID2 SeqID5	121 121	GCCGAATGGGTAATGATGAATGTAGGTTGGCCCCGAGAAGGAACTTTCACCATTGACAAT	
SeqID2 SeqID5	181 181	ATTTCACAGGTCGAGGAGAATCTTCGCCCCGGGGCCATATGGACACCCAGATCAAATC ""CAGC"""""G""""""""""""""""""""""""""	
SeqID2 SeqID5	241 241	CCTTATATTACCACGTGGAGATCCCTAGCCACAGACCCCCCTCCATGGGTTCGCCCATTC ""C""C""C""C""""GAG"""GAG"""G"""""C"""""C"""""GA""""GA""""""""	
SeqID2 SeqID5	301 301	CTACCCCTCCTAAGCATCCCAGGACAGATCCTCCCGAGCCTCTTTCGCCGCAACCTCTT ""G""""""C""C""""""C""C""C""C""""""""C""GAGC""C""G""C""G	
SeqID2 SeqID5	361 361	GCGCCGCAACCC_TC_TTCCCCCCA_CCCCGTCCTCTACCCCGTTCTCCCCAAACCAGAC ""C""C""G"""AG"GCC""""""T"AG"AG"""G""""""""G""G""""""""""	
SeqID2 SeqID5	418 421	CCCCCCAAGGCGCCTGTATTACCACCCAATCCTTCTTCCCCTTTAATTGATCTCTTAACA	
SeqID2 SeqID5	478 481	GAAGAGCCACCTCCCTATCCTGGGGGTCACGGGCCAACACCGCCGTCAGGCCCTAGAACC ""G"""""C""C""C""C""C""C""C""C""C""C""CAGC"""""C""G"""	
SeqID2 SeqID5	538 541	CCAACTGCCTCCCCGATTGCCATCCGGCTGCGAGAACGACGAGAAAATCCAGCTGAGAAA ""C""C"""AG"""C""C""""G"A"""""G""GA"GA"G""G""C""C""C"""""""G"	
SeqID2 SeqID5	598 601	TCTCAAGCCCTCCCCTTAAGGGAAGACCCCAAACAACAGACCCCAGTACTGGCCATTCTCG AGC""G"""""G"""C"G"""""G""""C""""AGC	
SeqID2 SeqID5	658 661	GCCTCTGACCTGTACAATTGGAAATTGCATAA_CCCCCCTTTCTCCCAGGACCCAGTGGC	
SeqID2 SeqID5	717 720	CCTAACTAACCTAATTGAGTCCATTTTAGTGACACATCAGCCAACCTGGGACGACTGCCA	
SeqID2 SeqID5	777 780	ACAGCTCTTACAGGCTCTCCTGACGGCAGAGAGAGAGACAAAGGGTCCTCCTTGAAGCCCGG"""""GC"G"""""C""C"""""""G""G""G""	
SeqID2 SeqID5	837 840	AAAGCAAGTTCCAGGCGAGGACGGACGGCCAACCCAGCTGCCCAATGTCGTTGACGAGGC	
SeqID2 SeqID5	897 900	TTTCCCCTTGACCCGTCCCAACTGGGATTTTTGTACGCCGGCAGGTAGGGAGCACCTACG	
SeqID2 SeqID5	957 960	CCTTTATCGCCAGTTGCTGTTAGCGGGGCTCCGCGGGGCTGCAAGACGCCCCACTAATTT G""G""CA"G"""C"""""C"G""C""GA"G""C""C""GA"G"""""C""C""	
SeqID2 SeqID5	1017 1020	nnnGnnnnnGnnnannGnnGnnGnnCnnGnnnnnGnnAnnCnnnAGCnnnCnGnnGnn	
SeqID2 SeqID5	1077 1080	ATTAAAAGAGGCTTACAGAATGTATACTCCCTATGACCCTGAGGACCCAGGGCAGGCTGC GC"G""G"""""C"""""C"""""C"""""C"""""C""""""	
SeqID2 SeqID5	1137 1140	TAGTGTTATCCTGTCCTTTATCTACCAGTCTAGCCCGGACATAAGAAATAAGTTACAAAG C""C""G""""AG"""C"""""AG"""C""""C"	

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SeqID2 SeqID5	1197 1200	GCTAGAAGGCCTACAGGGGTTCACACTGTCTGATTTGCTAAAAGAGGCAGAAAAGATATA
SeqID2 SeqID5	1257 1260	CAACAAAAGGGAAACCCCAGAGGAAAGGGAAGAAAGATTATGGCAGCGGCAGGAAGAAAG ####################
SeqID2 SeqID5	1317 1320	AGATAAAAAGCGCCATAAGGAGATGACTAAAGTTCTGGCCACAGTAGTTGCTCAGAATAGG""C""G""A"G""C"""""""""C""G""G""G""""""""
SeqID2 SeqID5	1377 1380	AGATAAGGATAGAGGGGAAAGTAAACTGGGAGATCAAAGGAAAATACCTCTGGGGAAAGA G""C"""""C""G""C""G""C""G""C""G""C""G"""C""C
SeqID2 SeqID5	1437 1440	
SeqID2 SeqID5	1497 1500	

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<u>Fig. 5</u>	DI ag	NA sequence comparison of the wild type "env" (gp70 region from Seq.ID1) painst the codon- and signal optimized "env" gene (gp70; Seq.ID8).
SeqID11 SeqID8	1 1	ATGGAAAGTCCAACGCACCCAAAACCCTCTAAAGATAAGACTCTCTCGTGGAACTTAGCG
SeqID11 SeqID8	61 61	TTTCTGGTGGGGATCTTATTTACAATAGACATAGGAATGGCCAATCCTAGTCCACACCAA
SeqID11 SeqID8	121 121	ATATATAATGTAACTTGGGTAATAACCAATGTACAAACTAACACCCAAGCTAACGCCACC
SeqID11 SeqID8	181 181	TCTATGTTAGGAACCTTAACCGATGCCTACCCTACCCTA
SeqID11 SeqID8	241 241	GTGGGAGACACCTGGGAACCTATAGTCCTAAACCCAACCAA
SeqID11 SeqID8	301 301	TACTCCTCCTCAAAATATGGATGTAAAACTACAGATAGAAAAAACAGCAACAGACATAC
SeqID11 SeqID8	361 361	CCCTTTTACGTCTGCCCCGGACATGCCCCCTCGTTGGGGCCAAAGGGAACACATTGTGGA
SeqID11 SeqID8	421 421	GGGGCACAAGATGGGTTTTGTGCCGCATGGGGATGTGAGACCACCGGAGAAGCTTGGTGG
SeqID11 SeqID8	481 481	AAGCCCACCTCCTCATGGGACTATATCACAGTAAAAAGAGGGAGTAGTCAGGACAATAGC
SeqID11 SeqID8	541 541	
SeqID11 SeqID8	601 601	TGGGACGGACCTAAGATGTGGGGATTGCGACTATACCGTACAGGATATGACCCTATCGCT
SeqID11 SeqID8	661 661	TTATTCACGGTGTCCCGGCAGGTATCAACCATTACGCCGCCTCAGGCAATGGGACCAAAC
SeqID11 SeqID8	721 721	CTAGTCTTACCTGATCAAAAACCCCCCATCCCGACAATCTCAAACAGGGTCCAAAGTGGCG
SeqID11 SeqID8	781 781	ACCCAGAGGCCCCAAACGAATGAAAGCGCCCCAAGGTCTGTTGCCCCCACCACCATGGGT
SeqID11 SeqID8	841 841	CCCAAACGGATTGGGACCGGAGATAGGTTAATAAATTTAGTACAAGGGACATACCTAGCC
SeqID11 SeqID8	901 901	TTAAATGCCACCGACCCCAACAAAACTAAAGACTGTTGGCTCTGCCTGGTTTCTCGACCA
SeqID11 SeqID8	961 961	CCCTATTACGAAGGGATTGCAATCTTAGGTAACTACAGCAACCAAACAAA
SeqID11 SeqID8	1021 1021	TCCTGCCTATCTACTCCGCAACACAAACTAACTATATCTGAAGTATCAGGGCAAGGAATG
SeqID11 SeqID8	1081 1081	TGCATAGGGACTGTTCCTAAAACCCACCAGGCTTTGTGCAATAAGACACAACAGGGACAT
SeqID11 SeqID8	1141 1141	ACAGGGCGCACTATCTAGCCGCCCCCAACGGCACCTATTGGGCCTGTAACACTGGACTC

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SeqID11 SeqID8	1201 1201	ACCCCATGCATTTCCATGGCGGTGCTCAATTGGACCTCTGATTTTTGTGTCTTAATCGAA
SeqID11 SeqID8	1261 1261	TTATGGCCCAGAGTGACTTACCATCAACCCGAATATGTGTACACACATTTTGCCAAAGCTC"G""""""""""
SeqID11 SeqID8	1321 1321	GTCAGGTTCCGAAGAGAACCAATATCACTAACGGTTGCCCTTATGTTGGGAGGACTTACT ""G"""""A"G"""A""G"""G"""C""C""G"""A""G""""G"""C"""G"""G
SeqID11 SeqID8	1381 1381	GTAGGGGCATAGCCGCGGGGTCGGAACAGGGACTAAAGCCCTCCTTGAAACAGCCTGA

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Fig. 6 DNA sequence comparison of the wild type "env" gene (Seq.ID1) against the codonand signal optimized "env" gene (gp85) (Seq.ID7).

SeqID1 SeqID7	1 1	ATGGAAAGTCCAACGCACCCAAAACCCTCTAAAGATAAGACTCTCTCGTGGAACTTAGCG """"GTCC""C""C"""""C""G"""""C""G""C"""""C""G""C""""""
SeqID1		TTTCTGGTGGGGATCTTATTTACAATAGACATAGGAATGGCCAATCCTAGTCCACACCAA
SeqID7	61	
SeqID1	121	ATATATATGTAACTTGGGTAATAACCAATGTACAAACTAACACCCAAGCTAACGCCACC
SeqID7	121	""C", "C", "", ", ", ", ", ", ", ", ", ", ", ",
SeqID1	181	TCTATGTTAGGAACCTTAACCGATGCCTACCCTACCTACATGTTGACTTATGTGACCTA
SeqID7	181	пиппинСпСпипСпипСпСпининаДпиппиСпипппСпипппСпиппппСпСпиппиппп
SeqID1	241	GTGGGAGACACCTGGGAACCTATAGTCCTAAACCCAACCAA
SeqID7	241	nnuuGnanuunaangaaCuaTCCGaaGaaaaaCaanaaaaaaaaaaaaaaaaaaaaaaa
CoaTD1	201	TACTCCTCCTCAAAATATGGATGTAAAACTACAGATAGAAAAAAAA
SeqID1		ununununun Caudaunnu CauCauGauGaunun CandauGauGaugaunuu Gaanau Caaa
SeqID7	301	
SeqID1	361	$\tt CCCTTTTACGTCTGCCCCGGACATGCCCCCTCGTTGGGGCCCAAAGGGAACACATTGTGGA$
SeqID7	361	ununCuuInuGununIunCunuunununuCCununCuuCuununCuuCuuCuununG
CoaTD1	421	${\tt GGGGCACAAGATGGGTTTTGTGCCGCATGGGGATGTGAGACCACCGGAGAAGCTTGGTGG}$
SeqID1	421	nnunCunGunauaCununuaLunCununuCununuAnnunuAnnunuHuncunGunCununun
SeqID7		
SeqID1	481	AAGCCCACCTCCTCATGGGACTATATCACAGTAAAAAGAGGGAGTAGTCAGGACAATAGC
SeqID7	481	пиниппппппппппппппппппппппппппппппппппп
SeqID1	541	TGTGAGGGAAAATGCAACCCCTGGTTTTGCAGTTCACCCAGAAGGGAAGACAAGCCTCT
SeqID7	541	unnunnuCunGunuunnunuunnuGCunuunnunuunuunuunuuuCunGuunuu
beq1D,	311	
SeqID1	601	${\tt TGGGACGGACCTAAGATGTGGGGATTGCGACTATACCGTACAGGATATGACCCTATCGCT}$
	601	nunnaLuncancanunnunnunununccanya.dundununya.dununucunnununununcuntunc
SeqID7		
SeqID1	661	TTATTCACGGTGTCCCGGCAGGTATCAACCATTACGCCGCCTCAGGCAATGGGACCAAAC
SeqID7	661	$C_{n}G_{n+n+1}M_{n+n+n+1}M_{n+n+n+1}G_{n+1}G_{n+1}G_{n+n+1}G_{n+1}G_{n+n+1}G_{n+n+1}G_{n+n+1}G_{n+n+1}G_{n+n+1}G_{n+n+1}G_{n+1}G_{n+n+1}G_{n+n+1}G_{n+n+1}G_{n+1}G_{n+1}G_{n+1}G_{n+1}G_{n+1}G$
CowID1	721	CTAGTCTTACCTGATCAAAAACCCCCCATCCCGACAATCTCAAACAGGGTCCAAAGTGGCG
SeqID1	721	""G""GC"G""""""C""G""G"""C"""A"G""G""G""G""G""G""""""C"""G"""G
SeqID7	/21	
SeqID1	781	
SeqID7	781	пинининининин вин Синини в ТСТ ини пи Сининин пи п Сининининин пи пи п
SeqID1	841	CCCAAACGGATTGGGACCGGAGATAGGTTAATAAATTTAGTACAAGGGACATACCTAGCC
SeqID7	841	
seq157	*	
SeqID1	901	TTAAATGCCACCGACCCCAACAAAACTAAAGACTGTTGGCTCTGCCTGGTTTCTCGACCA
SeqID7	901	$C_{11}G_{11}\ldots G_{11}\ldots G_{11}$
SeqID1	061	CCCTATTACGAAGGGATTGCAATCTTAGGTAACTACAGCAACCAAACAAA
_	961	
SeqID7		_
SeqID1	1021	TCCTGCCTATCTACTCCGCAACACAAACTAACTATATCTGAAGTATCAGGGCAAGGAAT
SeqID7	1020	CuuunnuuGuuCuuCuuCuuGuunuuGuuGuuCuuCuunuuGuuGuuTuuCuuGuuCuu
SeqID1	1080	GTGCATAGGGACTGTTCCTAAAACCCCACCAGGCTTTGTGCAATAAGACACAACAGGGACA
SeqID7	1080	$ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\$
		ma ca cocococoa cima mema cococococo a cococa coma minero de concena a cinco a cim
SeqID1	1140	TACAGGGGCGCACTATCTAGCCGCCCCCAACGGCACCTATTGGGCCTGTAACACTGGACT
SeqID7	1140	General and General General General and an anti-

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SeqID1 SeqID7	1200	CACCCCATGCATTTCCATGGCGGTGCTCAATTGGACCTCTGATTTTTGTGTCTTAATCGA G""""C""""C"""""G""C"""""G""C""""""C""C"
SeqID1 SeqID7	1260 1260	ATTATGGCCCAGAGTGACTTACCATCAACCCGAATATGTGTACACACATTTTGCCAAAGCGC"G""""""""""
SeqID1 SeqID7	1320 1320	TGTCAGGTTCCGAAGAGAACCAATATCACTAACGGTTGCCCTTATGTTGGGAGGACTTAC """G"""""""""""""""""""""""""""""""""
SeqID1 SeqID7	1380 1380	TGTAGGGGCATAGCCGCGGGGTCGGAACAGGGACTAAAGCCCTCCTTGAAACAGCCCA A""G"""""T""T""T""T"""""""""""""""""""
SeqID1 SeqID7	1440 1440	GTTCAGACAACTACAAATGGCCATGCACACAGACATCCAGGCCCTAGAAGAATCAATTAG
SeqID1 SeqID7	1500 1500	TGCCTTAGAAAAGTCCCTGACCTCCCTTTCTGAAGTAGTCTTACAAAACAGACGGGCCT C""T""""""A"""""A""""""A""""""""""A""""""
SeqID1 SeqID7	1560 1560	AGATATTCTATTCTTACAAGAGGGAGGGCTCTGTGCCGCATTGAAAGAAGAATGTTGCTT
SeqID1 SeqID7	1620 1620	CTATGCGGATCACACCGGACTCGTCCGAGACAATATGGCCAAATTAAGAGAAAGACTAAA
SeqID1 SeqID7	1680 1680	ACAGCGCAACAACTGTTTGACTCCCAACAGGGATGGTTTGAAGGATGGTTCAACAAGTC
SeqID1 SeqID7	1740 1740	
SeqID1 SeqID7	1800 1800	TCTCCTCTTCGGCCCATGCATCCTTAACCGATTAGTACAATTCGTAAAAGACAGAATATC
SeqID1 SeqID7	1860 1860	
SeqID1 SeqID7	1920 1920	

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<u>Fig. 7</u>	Protein sequence comparison of the wild type "gag" protein (Seq.ID4) against the protein sequence of the codon-optimized "gag" protein (Seq.ID6).
SeqID4 1 SeqID61	MGQTITTPLSLTLNHWSEVQARARNQGVEVRKKKWITLCEAEWVMMNVGWPREGTFTIDN
SeqID461 SeqID661	ISQVEERIFAPGPYGHPDQIPYITTWRSLATDPPPWVRPFLPPPKHPRTDPPEPLSPQPL
SeqID4121 SeqID6121	APQPSSPHPV_LYPVLPKPDPPKAPVLPPNPSSPLIDLLTEEPPPYPGGHGPTPPSGPRT
SeqID4180 SeqID6181	PTASPIAIRLRERRENPAEKSQALPLREDPNNRPQYWPFSASDLYNWKLHNPPFSQDPVA
SeqID4240 SeqID6241	LTNLIESILVTHQPTWDDCQQLLQALLTAEERQRVLLEARKQVPGEDGRPTQLPNVVDEA
SeqID4300 SeqID6301	FPLTRPNWDFCTPAGREHLRLYRQLLLAGLRGAARRPTNLAQVKQVVQGKEETPASFLER
SeqID4360 SeqID6361	LKEAYRMYTPYDPEDPGQAASVILSFIYQSSPDIRNKLQRLEGLQGFTLSDLLKEAEKIY
SeqID4420 SeqID6421	NKRETPEEREERLWQRQEERDKKRHKEMTKVLATVVAQNRDKDRGESKLGDQRKIPLGKD
SeqID4480 SeqID6481	QCAYCKEKGHWVRDCPKRPRKKPANSTLL

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Fig. 8 Protein sequence comparison of the wild type "env" protein (Seq.ID3) against the protein sequence of the codon- and signal optimized "env" protein (gp70) (Seq.ID10).

SeqID3 SeqID10	1 1	MESPTHPKPSKDKTLSWNLAFLVGILFTIDIGMANPSPHQIYNVTWVITNVQTNTQANAT
SeqID3 SeqID10	61 61	SMLGTLTDAYPTLHVDLCDLVGDTWEPIVLNPTNVKHGARYSSSKYGCKTTDRKKQQQTY
SeqID3 SeqID10		PFYVCPGHAPSLGPKGTHCGGAQDGFCAAWGCETTGEAWWKPTSSWDYITVKRGSSQDNS
SeqID3 SeqID10	181 181	CEGKCNPLVLQFTQKGRQASWDGPKMWGLRLYRTGYDPIALFTVSRQVSTITPPQAMGPN
SeqID3 SeqID10	241 241	LVLPDQKPPSRQSQTGSKVATQRPQTNESAPRSVAPTTMGPKRIGTGDRLINLVQGTYLA
SeqID3 SeqID10	301 301	LNATDPNKTKDCWLCLVSRPPYYEGIAILGNYSNQTNPPPSCLSTPQHKLTISEVSGQGM
SeqID3 SeqID10		CIGTVPKTHQALCNKTQQGHTGAHYLAAPNGTYWACNTGLTPCISMAVLNWTSDFCVLIE
SeqID3 SeqID10	421 421	LWPRVTYHQPEYVYTHFAKAVRFRREPISLTVALMLGGLTVGGIAAGVGTGTKALLETA

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Fig. 9	t	Protein sequence comparison of the wild type "env" protein (Seq.ID3) against the protein sequence of the codon- and signal optimized "env" protein (gp85) Seq.ID9).
SeqID3 SeqID9	1	MESPTHPKPSKDKTLSWNLAFLVGILFTIDIGMANPSPHQIYNVTWVITNVQTNTQANAT
SeqID3 SeqID9	61 61	SMLGTLTDAYPTLHVDLCDLVGDTWEPIVLNPTNVKHGARYSSKYGCKTTDRKKQQQTY
SeqID3 SeqID9	121 121	PFYVCPGHAPSLGPKGTHCGGAQDGFCAAWGCETTGEAWWKPTSSWDYITVKRGSSQDNS
SeqID3 SeqID9	181 181	CEGKCNPLVLQFTQKGRQASWDGPKMWGLRLYRTGYDPIALFTVSRQVSTITPPQAMGPN
SeqID3 SeqID9	241 241	LVLPDQKPPSRQSQTGSKVATQRPQTNESAPRSVAPTTMGPKRIGTGDRLINLVQGTYLA
SeqID3 SeqID9	301 301	LNATDPNKTKDCWLCLVSRPPYYEGIAILGNYSNQTNPPPSCLSTPQHKLTISEVSGQGM
SeqID3 SeqID9	361 361	CIGTVPKTHQALCNKTQQGHTGAHYLAAPNGTYWACNTGLTPCISMAVLNWTSDFCVLIE
SeqID3 SeqID9	421 421	LWPRVTYHQPEYVYTHFAKAVRFRREPISLTVALMLGGLTVGGIAAGVGTGTKALLETAQ
SeqID3 SeqID9	481 481	FRQLQMAMHTDIQALEESISALEKSLTSLSEVVLQNRRGLDILFLQEGGLCAALKEECCF
SeqID3 SeqID9	541 541	YADHTGLVRDNMAKLRERLKQRQQLFDSQQGWFEGWFNKSPWFTTLISSIMGPLLILLLI
SeqID3 SeqID9	601 601	LLFGPCILNRLVQFVKDRISVVQALILTQQYQQIKQYDPDRP